

GenCore version 5.1.6
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OM protein - protein search, using BW model

Run on: September 16, 2005, 08:30:51 ; Search time 30.5 Seconds
(without alignments)
974.109 Million cell updates/sec

Title: US-10-643-627-6

Perfect score: 398
Sequence: 1 MNVLSEGTSTVAETFTISVM.....KSRKSSSYSSSTTTTKTSY 398

Scoring table: OLIGO
Gapop 60.0 , Gapext 60.0

Searched: 513545 seqs, 74649064 residues

Word size : 0

Total number of hits satisfying chosen parameters: 513545

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: listing first 1000 summaries

Database :

Issued Patents:AA:*
1: /cgn2_6/ptodata/1/1aa/5A.COMB.pep:*
2: /cgn2_6/ptodata/1/1aa/5B.COMB.pep:*
3: /cgn2_6/ptodata/1/1aa/6A.COMB.pep:*
4: /cgn2_6/ptodata/1/1aa/6B.COMB.pep:*
5: /cgn2_6/ptodata/1/1aa/6C.COMB.pep:*
6: /cgn2_6/ptodata/1/1aa/6D.COMB.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match length	ID	Description
1	398	100.0	398 3 US-08-486-673B-6	Sequence 6, Appli
2	291	73.1	398 1 US-08-097-938-6	Sequence 6, Appli
3	291	73.1	398 1 US-08-476-000-6	Sequence 6, Appli
4	291	73.1	398 1 US-08-472-840-6	Sequence 6, Appli
5	291	73.1	398 2 US-08-476-976-6	Sequence 6, Appli
6	291	73.1	398 3 US-08-474-410-6	Sequence 6, Appli
7	217	54.5	398 1 US-08-097-938-4	Sequence 4, Appli
8	217	54.5	398 1 US-08-476-000-4	Sequence 4, Appli
9	217	54.5	398 1 US-08-472-840-4	Sequence 4, Appli
10	217	54.5	398 2 US-08-476-976-4	Sequence 4, Appli
11	217	54.5	398 3 US-08-474-410-4	Sequence 4, Appli
12	217	54.5	398 3 US-08-486-673B-4	Sequence 4, Appli
13	155	28.6	394 3 US-08-742-440A-8	Sequence 8, Appli
14	114	28.6	397 3 US-08-486-673B-63	Sequence 63, Appli
15	91	22.9	397 1 US-08-476-000-63	Sequence 63, Appli
16	91	22.9	397 1 US-08-472-840-63	Sequence 63, Appli
17	91	22.9	397 2 US-08-476-976-63	Sequence 63, Appli
18	91	22.9	397 3 US-08-474-410-63	Sequence 63, Appli
19	34	8.5	395 1 US-08-097-938-2	Sequence 2, Appli
20	34	8.5	395 1 US-08-097-938-5	Sequence 5, Appli
21	34	8.5	395 1 US-08-476-000-2	Sequence 2, Appli
22	34	8.5	395 1 US-08-476-000-5	Sequence 5, Appli
23	34	8.5	395 1 US-08-472-840-2	Sequence 2, Appli
24	34	8.5	395 1 US-08-472-840-5	Sequence 5, Appli
25	34	8.5	395 2 US-08-476-976-2	Sequence 2, Appli
26	34	8.5	395 2 US-08-476-976-5	Sequence 5, Appli
27	34	8.5	395 3 US-08-474-410-2	Sequence 2, Appli

28	34	8.5	395 3 US-08-474-410-5	Sequence 5, Appli
29	34	8.5	395 3 US-08-486-673B-2	Sequence 2, Appli
30	34	8.5	395 3 US-08-486-673B-5	Sequence 5, Appli
31	34	8.5	399 1 US-08-476-000-61	Sequence 61, Appli
32	34	8.5	399 1 US-08-472-840-61	Sequence 61, Appli
33	34	8.5	399 2 US-08-476-976-61	Sequence 61, Appli
34	34	8.5	399 3 US-08-474-410-61	Sequence 61, Appli
35	34	8.5	399 3 US-08-486-673B-61	Sequence 61, Appli
36	34	8.5	399 3 US-08-486-673B-61	Sequence 61, Appli
37	36	3.8	25 4 US-09-472-130A-19	Sequence 19, Appli
38	36	3.8	31 1 US-08-097-938-21	Sequence 21, Appli
39	36	3.8	31 1 US-08-476-000-21	Sequence 21, Appli
40	36	3.8	31 2 US-08-476-976-21	Sequence 21, Appli
41	36	3.8	31 3 US-08-474-410-21	Sequence 21, Appli
42	43	3.8	31 3 US-08-486-673B-21	Sequence 21, Appli
43	43	3.8	13 1 US-08-097-938-41	Sequence 41, Appli
44	43	3.8	13 1 US-08-476-000-41	Sequence 41, Appli
45	43	3.8	13 1 US-08-472-840-41	Sequence 41, Appli
46	43	3.8	13 2 US-08-476-976-41	Sequence 41, Appli
47	43	3.8	13 3 US-08-474-410-41	Sequence 41, Appli
48	43	3.8	13 3 US-08-486-673B-41	Sequence 41, Appli
49	43	3.8	13 3 US-08-486-673B-41	Sequence 41, Appli
50	51	2.5	26 1 US-07-789-184-217	Sequence 217, Appli
51	51	2.5	26 1 US-07-789-184-217	Sequence 217, Appli
52	51	2.5	26 1 US-07-789-184-217	Sequence 217, Appli
53	51	2.5	26 1 US-07-789-184-217	Sequence 217, Appli
54	51	2.5	26 1 US-08-475-263-214	Sequence 214, Appli
55	51	2.5	26 1 US-08-475-263-214	Sequence 214, Appli
56	51	2.5	26 1 US-08-485-886-103	Sequence 103, Appli
57	51	2.5	26 1 US-08-485-886-214	Sequence 214, Appli
58	51	2.5	26 1 US-08-485-886-217	Sequence 217, Appli
59	51	2.5	26 2 US-08-477-362-103	Sequence 103, Appli
60	51	2.5	26 2 US-08-477-362-214	Sequence 214, Appli
61	61	2.5	26 2 US-08-477-362-217	Sequence 217, Appli
62	61	2.5	26 2 US-08-477-134-103	Sequence 103, Appli
63	61	2.5	26 2 US-08-477-134-214	Sequence 214, Appli
64	61	2.5	26 2 US-08-477-134-217	Sequence 217, Appli
65	61	2.5	26 3 US-08-473-489A-103	Sequence 103, Appli
66	61	2.5	26 3 US-08-473-489A-214	Sequence 214, Appli
67	61	2.5	26 3 US-08-473-489A-217	Sequence 217, Appli
68	61	2.5	26 3 US-08-485-695-103	Sequence 103, Appli
69	61	2.5	26 3 US-08-485-695-214	Sequence 214, Appli
70	61	2.5	26 3 US-08-485-695-217	Sequence 217, Appli
71	71	2.5	26 3 US-08-018-760-103	Sequence 103, Appli
72	71	2.5	26 3 US-08-018-760-214	Sequence 214, Appli
73	71	2.5	26 3 US-08-018-760-217	Sequence 217, Appli
74	74	2.5	408 2 US-08-742-440A-6	Sequence 6, Appli
75	74	2.5	425 1 US-07-657-769B-69	Sequence 69, Appli
76	74	2.5	425 1 US-08-097-938-7	Sequence 7, Appli
77	74	2.5	425 1 US-08-313-553-13	Sequence 13, Appli
78	74	2.5	425 1 US-07-789-184-220	Sequence 220, Appli
79	74	2.5	425 1 US-08-476-000-7	Sequence 7, Appli
80	74	2.5	425 1 US-08-476-263-220	Sequence 220, Appli
81	74	2.5	425 1 US-08-472-840-7	Sequence 7, Appli
82	74	2.5	425 1 US-08-485-886-220	Sequence 220, Appli
83	74	2.5	425 2 US-08-477-362-220	Sequence 220, Appli
84	74	2.5	425 2 US-08-477-134-220	Sequence 220, Appli
85	74	2.5	425 2 US-08-911-320A-3	Sequence 3, Appli
86	74	2.5	425 2 US-08-476-976-7	Sequence 7, Appli
87	74	2.5	425 2 US-08-742-440A-7	Sequence 7, Appli
88	74	2.5	425 2 US-08-560-098A-57	Sequence 57, Appli
89	74	2.5	425 3 US-08-767-993-13	Sequence 13, Appli
90	74	2.5	425 3 US-08-473-489A-220	Sequence 220, Appli
91	74	2.5	425 3 US-08-474-410-7	Sequence 7, Appli
92	74	2.5	425 3 US-08-485-695-220	Sequence 220, Appli
93	74	2.5	425 3 US-08-217-101-3	Sequence 217, Appli
94	74	2.5	425 3 US-08-018-760-220	Sequence 220, Appli
95	74	2.5	425 3 US-08-486-673B-7	Sequence 7, Appli
96	74	2.5	425 4 US-09-054-272-53	Sequence 53, Appli
97	74	2.5	12 1 US-08-097-938-42	Sequence 42, Appli
98	74	2.5	12 1 US-08-476-000-42	Sequence 42, Appli
99	74	2.5	12 1 US-08-472-840-42	Sequence 42, Appli
100	74	2.5	12 2 US-08-476-976-42	Sequence 42, Appli

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OM protein - protein search, using sw model

Run on: September 16, 2005, 08:17:59 ; Search time 119.5 Seconds
(without alignments)
1288.122 Million cell updates/sec

Title: US-10-643-627-6

Perfect score: 398

Sequence: 1 MNVLSFRQTSVTAETPISWM.....KHSRKSSTSSYSTTVKTSY 398

Scoring table: OLIGO

Gapop 60.0 , Gapext 60.0

Searched: 2105692 seqs, 386760381 residues

Word size: 0

Total number of hits satisfying chosen parameters: 2105692

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Listing first 1000 summaries

Database:

1: Genesegp16Dec04:*
2: Genesegp1990s:*
3: Genesegp2000s:*
4: Genesegp2001s:*
5: Genesegp2002s:*
6: Genesegp2003as:*
7: Genesegp2003bs:*
8: Genesegp2004s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	217	54.5	397	3 AAB35641	AAB35641 Human PAR
2	217	54.5	397	5 AAB26678	AAB26678 Human coa
3	217	54.5	397	6 ABG73508	ABG73508 Human par
4	217	54.5	397	7 ADG62812	ADG62812 Human Pro
5	217	54.5	397	8 ADI61221	ADI61221 Human coa
6	217	54.5	397	8 ADG29311	ADG29311 Human GPC
7	217	54.5	397	8 ADG74020	ADG74020 Human G-P
8	217	54.5	398	2 AAB66921	AAB66921 Human C14
9	217	54.5	398	2 AAW01953	AAW01953 Human C14
10	162	40.7	320	8 ADI28655	ADI28655 Human mod
11	162	40.7	341	8 ADI28654	ADI28654 Human mod
12	162	40.7	355	8 ADI28653	ADI28653 Human mod
13	162	40.7	397	6 ABP81907	ABP81907 Human pro
14	162	40.7	397	7 ADK32594	ADK32594 Hematolog
15	162	40.7	397	7 ADN39997	ADN39997 Cancer/an
16	162	40.7	397	8 ADK46675	ADK46675 Cancer-as
17	155	38.9	394	2 AAW51408	AAW51408 Human pro
18	155	38.9	397	2 AAB66923	AAB66923 Human C14
19	91	22.9	397	2 AAW01955	AAW01955 Human C14
20	57	14.3	389	8 ADO28601	ADO28601 Human PAR
21	37	9.3	397	7 ADG62810	ADG62810 Rat Prote
22	35	8.8	58	5 ABU67239	ABU67239 G-protein
23	35	8.8	58	6 ABP54020	ABP54020 Human pro
24	35	8.8	58	6 ADO28760	ADO28760 Human pro
25	35	8.8	58	8 ADO05264	ADO05264 Proteinas

26	34	8.5	395	2 AAB66920	AAB66920 Murine C1
27	34	8.5	395	2 AAW01952	AAW01952 Murine C1
28	34	8.5	399	2 AAB66922	AAB66922 Murine C1
29	34	8.5	399	2 AAW01954	AAW01954 Murine C1
30	34	8.5	399	7 ABR63562	ABR63562 Delayed h
31	34	8.5	399	7 ADO29312	ADO29312 Mouse GPC
32	25	6.3	54	8 ADI28666	ADI28666 Human pro
33	22	5.5	25	3 AAY45037	AAY45037 Human pro
34	22	5.5	21	5 AAW04020	AAW04020 PAR2 pepd
35	20	5.0	21	5 AAW04021	AAW04021 PAR2 pepd
36	19	4.8	20	6 ABP82705	ABP82705 G protein
37	18	4.5	18	6 ABP82706	ABP82706 G protein
38	18	4.5	18	6 ABP82707	ABP82707 G protein
39	15	3.8	68	5 ADK35286	ADK35286 Novel hum
40	14	3.5	20	3 AAB35651	AAB35651 Human PAR
41	12	3.0	12	2 AAW76405	AAW76405 Human PAR
42	12	3.0	16	6 ABP82708	ABP82708 G protein
43	11	2.8	20	3 AAB35652	AAB35652 Mouse PAR
44	10	2.5	13	2 AAB66890	AAB66890 Agonist P
45	10	2.5	13	2 AAW01923	AAW01923 C14o rece
46	10	2.5	15	5 ABG35288	ABG35288 Human PAR
47	10	2.5	15	5 ABG35287	ABG35287 Human PAR
48	10	2.5	15	5 ABG35289	ABG35289 Human PAR
49	10	2.5	15	5 ABG35285	ABG35285 Human PAR
50	10	2.5	26	2 AAR27238	AAR27238 Thrombin
51	10	2.5	319	4 AAB82760	AAB82760 Rat G-pro
52	10	2.5	319	8 ADO29401	ADO29401 Mouse GPC
53	10	2.5	371	5 ABG35299	ABG35299 Human PAR
54	10	2.5	374	2 AAW51406	AAW51406 Human pro
55	10	2.5	374	6 ABG73509	ABG73509 Human par
56	10	2.5	374	6 ABP81908	ABP81908 Human pro
57	10	2.5	374	7 ADG67661	ADG67661 Human P2R
58	10	2.5	374	8 ADO29313	ADO29313 Human GPC
59	10	2.5	374	8 ADQ97469	ADQ97469 Human can
60	10	2.5	374	8 ADQ39889	ADQ39889 Human myo
61	10	2.5	374	8 ADG32990	ADG32990 Proteinas
62	10	2.5	402	5 ABG35298	ABG35298 Human PAR
63	10	2.5	425	5 AAR27240	AAR27240 Human thr
64	10	2.5	425	5 AAB60698	AAB60698 Fragment
65	10	2.5	425	2 AAW51407	AAW51407 Human pro
66	10	2.5	425	2 AAY49570	AAY49570 Human thr
67	10	2.5	425	2 AAE17032	AAE17032 Human thr
68	10	2.5	425	5 ABG35300	ABG35300 Human PAR
69	10	2.5	425	5 AAG80697	AAG80697 Human thr
70	10	2.5	425	6 ABG73511	ABG73511 Human thr
71	10	2.5	425	6 ABR47449	ABR47449 Breast ca
72	10	2.5	425	6 ABP81919	ABP81919 Human thr
73	10	2.5	425	7 ADE58075	ADE58075 Human pro
74	10	2.5	425	7 ADE58071	ADE58071 Human pro
75	10	2.5	425	7 ADG89876	ADG89876 Human coa
76	10	2.5	425	8 ADL14208	ADL14208 Novel hum
77	10	2.5	425	8 ADN04016	ADN04016 Antipsoit
78	10	2.5	425	8 ADO29309	ADO29309 Human GPC
79	10	2.5	425	8 ADQ18985	ADQ18985 Human sof
80	10	2.5	425	8 ADI45608	ADI45608 Human G-P
81	10	2.5	425	8 ADS84489	ADS84489 Human pro
82	10	2.5	426	3 AAY45035	AAY45035 Human thr
83	10	2.5	430	8 ADO29310	ADO29310 Mouse GPC
84	10	2.5	432	7 ADE58073	ADE58073 Rat Prote
85	10	2.5	432	7 ADE58069	ADE58069 Rat Prote
86	10	2.5	892	2 AAW16314	AAW16314 Human thr
87	10	2.5	12	2 AAB66891	AAB66891 Agonist P
88	9	2.3	12	2 AAW01924	AAW01924 C14o rece
89	9	2.3	15	5 ABG35284	ABG35284 Human PAR
90	9	2.3	17	2 AAW76405	AAW76405 Human PAR
91	9	2.3	137	4 AAW41295	AAW41295 Propionib
92	9	2.3	137	4 AAW76404	AAW76404 Rat PAR-2
93	9	2.3	137	4 AAW76404	AAW76404 Rat PAR-2
94	8	2.0	8	2 AAR66892	AAR66892 Agonist P
95	8	2.0	11	2 AAW01925	AAW01925 C14o rece
96	8	2.0	15	5 ABG35283	ABG35283 Human PAR
97	8	2.0	20	2 AAW76404	AAW76404 Rat PAR-2
98	8	2.0	20	2 AAW76404	AAW76404 Rat PAR-2

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OM protein - protein search, using sw model

Run on: September 16, 2005, 08:17:59 / Search time 119.5 Seconds
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1288.122 Million cell updates/sec

Title: US-10-643-627-4

Perfect score: 398

Sequence: 1 MNVLSPFQTSVTAETFSVM.....KHSRKSSYSSTSTVTKTSY 398

Scoring table: OLIGO

Gapop 60.0, Gapext 60.0

Searched: 2105692 seqs, 386760381 residues

Word size: 0

Total number of hits satisfying chosen parameters: 2105692

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Listing first 1000 summaries

Database: A_Geneseq_16Dec04:*

1: geneseq19808:*

2: geneseq19908:*

3: geneseq20008:*

4: geneseq20018:*

5: geneseq20028:*

6: geneseq20038:*

7: geneseq20048:*

8: geneseq20048:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	398	100.0	398	2	AAR66921	Aar66921 Human C14
2	398	100.0	398	2	AAW01953	Aaw01953 Human C14
3	370	93.0	397	3	AAE26678	Aae26678 Human PAR
4	370	93.0	397	5	AAE26678	Aae26678 Human PAR
5	370	93.0	397	6	ABG73508	Abg73508 Human pro
6	370	93.0	397	7	ADG62812	Adg62812 Human pro
7	370	93.0	397	8	ADG62812	Adg62812 Human pro
8	370	93.0	397	8	ADG62812	Adg62812 Human pro
9	367	92.2	397	6	ADL61221	Adl61221 Human pro
10	367	92.2	397	6	ADL61221	Adl61221 Human pro
11	269	67.6	397	7	ADK52594	Adk52594 Hematolog
12	269	67.6	397	7	ADK52594	Adk52594 Hematolog
13	269	67.6	397	7	ADK52594	Adk52594 Hematolog
14	259	65.1	320	8	AD128655	Ad128655 Human mod
15	259	65.1	320	8	AD128655	Ad128655 Human mod
16	259	65.1	320	8	AD128655	Ad128655 Human mod
17	211	53.0	394	2	AAW51408	Aaw51408 Human pro
18	91	22.9	397	2	AAW66923	Aaw66923 Human C14
19	91	22.9	397	2	AAW66923	Aaw66923 Human C14
20	58	14.6	58	5	ABU67239	Abu67239 G-protein
21	58	14.6	58	6	ABU67239	Abu67239 G-protein
22	58	14.6	58	6	ABU67239	Abu67239 G-protein
23	58	14.6	58	8	ADG28760	Adg28760 Human pro
24	57	14.3	389	8	ADG28760	Adg28760 Human pro
25	54	13.6	54	8	AD128666	Ad128666 Human pro

26	37	9.3	397	7	ADG62810	Adg62810 Rat Prote
27	34	8.5	395	2	AAR66920	Aar66920 Murine C1
28	34	8.5	395	2	AAW01952	Aaw01952 Murine C1
29	34	8.5	399	2	AAR66922	Aar66922 Murine C1
30	34	8.5	399	2	AAW01954	Aaw01954 Murine C1
31	34	8.5	399	2	ABR63562	AbR63562 Delayed h
32	34	8.5	399	8	ADG29312	AdG29312 Mouse GPC
33	25	6.3	25	3	AAV45037	Aav45037 Human pro
34	21	5.3	21	5	AAU10420	Aau10420 PAR2 pepd
35	20	5.0	21	5	AAU10421	Aau10421 PAR2 pepd
36	19	4.8	20	6	ABP82705	Abp82705 G protein
37	18	4.5	18	6	ABP82706	Abp82706 G protein
38	18	4.5	18	6	ABP82707	Abp82707 G protein
39	16	4.0	16	6	ABP82708	Abp82708 G protein
40	15	3.8	16	6	ABP82708	Abp82708 G protein
41	14	3.5	20	3	AAW65651	Aaw65651 Human PAR
42	12	3.0	12	2	AAW76406	Aaw76406 Mouse PAR
43	11	2.8	20	3	AAW65652	Aaw65652 Mouse PAR
44	11	2.5	13	2	AAW66890	Aaw66890 Agonist p
45	10	2.5	13	2	AAW66890	Aaw66890 Agonist p
46	10	2.5	15	5	ABG35288	Abg35288 Human PAR
47	10	2.5	15	5	ABG35286	Abg35286 Human PAR
48	10	2.5	15	5	ABG35287	Abg35287 Human PAR
49	10	2.5	15	5	ABG35289	Abg35289 Human PAR
50	10	2.5	15	5	ABG35285	Abg35285 Human PAR
51	10	2.5	26	2	AAW72328	Aaw72328 Thrombin
52	10	2.5	319	4	AAW82760	Aaw82760 Rat G-pro
53	10	2.5	319	4	ADG29401	Adg29401 Mouse GPC
54	10	2.5	371	5	ABG35299	Abg35299 Human PAR
55	10	2.5	374	2	AAW51406	Aaw51406 Human pro
56	10	2.5	374	6	ABG73509	Abg73509 Human par
57	10	2.5	374	6	ABP81908	Abp81908 Human pro
58	10	2.5	374	6	ADG67661	Adg67661 Human P2R
59	10	2.5	374	8	ADG29313	AdG29313 Human GPC
60	10	2.5	374	8	ADG29313	AdG29313 Human GPC
61	10	2.5	374	8	ADG29313	AdG29313 Human GPC
62	10	2.5	374	8	ADG29313	AdG29313 Human GPC
63	10	2.5	374	8	ADG29313	AdG29313 Human GPC
64	10	2.5	425	5	ABG35298	Abg35298 Human PAR
65	10	2.5	425	5	AAW7240	Aaw7240 Human thr
66	10	2.5	425	2	AAW60698	Aaw60698 Fragment
67	10	2.5	425	2	AAW51407	Aaw51407 Human pro
68	10	2.5	425	2	AAW49570	Aaw49570 Human thr
69	10	2.5	425	5	ABG35300	Abg35300 Human thr
70	10	2.5	425	5	AAW60697	Aaw60697 Human thr
71	10	2.5	425	6	ABG73511	Abg73511 Human thr
72	10	2.5	425	6	ABR47449	AbR47449 Breast ca
73	10	2.5	425	6	ABP81919	Abp81919 Human thr
74	10	2.5	425	7	ADG58075	Adg58075 Human pro
75	10	2.5	425	7	ADG58071	Adg58071 Human pro
76	10	2.5	425	7	ADG58071	Adg58071 Human pro
77	10	2.5	425	8	ADG58071	Adg58071 Human pro
78	10	2.5	425	8	ADL14208	Adl14208 Novel hum
79	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
80	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
81	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
82	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
83	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
84	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
85	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
86	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
87	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
88	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
89	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
90	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
91	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
92	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
93	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
94	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
95	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
96	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
97	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC
98	10	2.5	425	8	ADG29310	AdG29310 Mouse GPC

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